

## SEQUENCE LISTING

- <110> KNAUF, VIC C. KRIDL, JEAN C.
- <120> METHODS AND COMPOSITIONS FOR REGULATED TRANSCRIPTION AND EXPRESSION OF HETEROLOGOUS GENES
- <130> 16518.052
- <140> US 09/782,130
- <141> 2001-02-12
- <150> US 09/232,861
- <151> 1999-01-15
- <150> US 08/812,665
- <151> 1997-03-07
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- <151> 1995-06-07
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- <151> 1993-08-10
- <150> US 07/526,123
- <151> 1990-05-21
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- <150> US 06/692,605
- <151> 1985-01-17
- <150> US 07/582,241
- <151> 1990-09-14
- <150> US 07/188,361
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- <151> 1988-01-25
- <150> US 07/078,538
- <151> 1987-07-28

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		_					_							tgg Trp		964
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	tgt Cys												tgaa	aaaa	atc	1061
ggti	caat	at o	ccggt	taaq	gc tt	taga	ataa	ato	gtgtg	gtgt	tggt	tata	at t	taaç	gactct	1121
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<211> 344

<212> PRT

<213> Brassica campestris

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- Tyr Ala Asp Ala Asn Glu Lys Asn Asn Arg Tyr Ala Val Phe Lys Arg 50 55 60
- Asn Val Glu Arg Ile Glu Arg Leu Asn Asp Val Gln Ser Gly Leu Thr
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- Phe Lys Leu Ala Val Asn Gln Phe Ala Asp Leu Thr Asn Glu Glu Phe 85 90 95
- Arg Ser Met Tyr Thr Gly Phe Lys Gly Asn Ser Val Leu Ser Ser Arg 100 105 110
- Thr Lys Pro Thr Ser Phe Arg Tyr Gln Asn Val Ser Ser Asp Ala Leu 115 120 125
- Pro Val Ser Val Asp Trp Arg Lys Lys Gly Ala Val Thr Pro Ile Lys 130 135 140
- Asp Gln Gly Leu Cys Gly Ser Cys Trp Ala Phe Ser Ala Val Ala Ala 145 150 155 160
- Ile Glu Gly Val Ala Gln Ile Lys Lys Gly Lys Leu Ile Ser Leu Ser 165 170 175
- Glu Gln Glu Leu Val Asp Cys Asp Thr Asn Asp Asp Gly Cys Met Gly
  180 185 190
- Gly Tyr Met Asn Ser Ala Phe Asn Tyr Thr Met Thr Thr Gly Gly Leu 195 200 205
- Thr Ser Glu Ser Asn Tyr Pro Tyr Lys Ser Thr Asp Gly Thr Cys Asn 210 215 220
- Phe Asn Lys Thr Lys Gln Ile Ala Thr Ser Ile Lys Gly Phe Glu Asp 225 230 235 240
- Val Pro Ala Asn Asp Glu Lys Ala Leu Met Lys Ala Val Ala His His 245 250 255
- Pro Val Ser Ile Gly Ile Ala Gly Gly Asp Ile Gly Phe Gln Phe Tyr 260 265 270
- Ser Ser Gly Val Phe Ser Gly Glu Cys Thr Thr His Leu Asp His Gly 275 280 285
- Val Thr Ala Val Gly Tyr Gly Arg Ser Lys Asn Gly Leu Lys Tyr Trp 290 295 300
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315

320

305

310

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atg a Met I	_		_				_	_		-	_				-	102
gat a Asp N																150
ccc o Pro I 40																198
ttg (																246
att a		Leu	_		Phe	Cys	_	Glu	Lys	Thr	Asp	_	Tyr	Gly		294
aca t		_		_		_	_		tgaa	acaat	at o	caatg	gatc	ta		341
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Lys Leu Leu Pro Thr Asn Ile Leu Gly Leu Cys Asn Glu Pro Cys Ser
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Ser Asn Ser Asp Cys Ile Gly Ile Thr Leu Cys Gln Phe Cys Lys Glu
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Lys Thr Asp Gln Tyr Gly Leu Thr Tyr Arg Thr Cys Asn Leu Leu Pro
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Ala

subunit peptide sequence

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gaaaacatgg atgtgaaaaa tactttttgt taaaagtaaa aaaaaatgtg aaattttgtt 180
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2377

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Lys Leu Leu Pro Thr Asn Ile Leu Gly Leu Cys Asn Glu Pro Cys Ser 50 55 60

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<sup>&</sup>lt;211> 96

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Lycopersicon esculentum

<sup>&</sup>lt;210> 26

<sup>&</sup>lt;211> 2207

<sup>&</sup>lt;212> DNA

<sup>&</sup>lt;213> Lycopersicon esculentum

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